PATENT COOPERATION TREATY

From the INTERNATIONAL SEARCHING AUTHORITY							
To:		PCT					
Ström & Gulliksson IP AB Box 7086 103 87 Stockholm Sverige	WRITTEN OPINION OF THE INTERNATIONAL SEARCHING AUTHORITY (PCT Rule 43 <i>bis.</i> 1)						
	Date of mailing (day/month/year)	2. 0 -0 1- 2005					
Applicant's or agent's file reference 2031002PC/ko	FOR FURTHER ACTION See paragraph 2 below						
International application No. International filing dat PCT/FI 2004/000540 15.09.2004	e (day/month/year)	Priority date (day/month/year) 15.09.2003					
International Patent Classification (IPC) or both national classification (IPC) at 15/70, C12N 15/61	cation and IPC						
Applicant Fit Biotech OYJ PLC et al							
1. This opinion contains indications relating to the following items: Box No. I Basis of the opinion							
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International application No.

PCT/FI 2004/000540

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		and	23.1(b)).									
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3.	In addition, in the case that more than one version or copy of a sequence listing and/or table relating thereto has been filed or furnished, the required statements that the information in the subsequent or additional copies is identical to that in the application as filed or does not go beyond the application as filed, as appropriate, were furnished.								s been al to			
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Box No. V Reasoned statement under Rule 43bis.1(a)(i) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement

1. Statement			
Novelty (N)	Claims	1-7, 9-24	YES
·	Claims	8	NO
Inventive step (IS)	Claims	1-7, 16-24	YES
Mysian or one (m)	Claims	8-15	NO
Industrial applicability (IA)	Claims	1-24	YES
mount of the same	Claims		NO

2. Citations and explanations:

The present application relates to an antibiotic resistance-free selection system, which is based on the use of an arab gene, a complementary sequence thereof or a catalytically active fragment thereof as a selection marker carried on a plasmid which is inserted in a bacterial strain deficient of the arab gene. Further described, are vectors comprising the arab gene, E. coli strains deficient of the arab gene and a method of selecting the cells transformed with a plasmid containing the arab gene.

The independent claims 1, 8, 13-22, are not considered to be sufficiently defined since they refer merely to the gene name "araD". In order to render the subject-matter of the claims clear and defined (PCT Art 6), the gene should be defined in the claims by reference to the sequence identity number(s) of the nucleotide sequence(s) of the gene, as disclosed in the sequence listing of the application.

Regarding the use of a vector carrying a complementary sequence of the araD gene (included in the subject-matter of claims 1, 8 and 22) it seems unlikely that the protein which is encoded by the complementary sequence of the araD gene would have the same functions as the protein encoded by the araD gene (unless the araD DNA sequence were a palindrome). Consequently, a selection system comprising a vector carrying the complementary sequence of the araD gene is not considered to function properly.

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Supplemental Box

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Reference will be made to the following documents cited in the international search report:

- D1) Protein Science, 4: 1648-1650 (1995), Andersson A et al.
- D2) W002090558 A1
- D3) Carcinogenesis, 14(2): 303-305 (1993), Ariza R R et al.

D1 discloses the vector pAA1 comprising the araD gene sequence. The E. coli strain JM109 is transformed with pAA1 in order to overexpress the araD gene and to purify the resulting enzyme, L-ribulose-5-phosphate 4-epimerase (see page 1648, column 2, paragraph 2-3 and the abstract).

Thus, the subject-matter of claim 8 lacks novelty. The fact that the araD gene will function as a selection marker when the vector is inserted in a selection system according to the invention does not confer novelty to the vector as such.

The vector according to dependent claims 9-12 may be novel. However, the construction of expression vectors is well known in the art. D2 discloses an expression vector, which lacks a papilloma virus origin of replication. The vector sequence encoding a nuclear-anchoring DNA comprises a protein operatively linked to a heterologous promoter, and a multimerized DNA sequence forming a binding site for the nuclear anchoring protein. Consequently, it is considered obvious to a person skilled in the art to combine the teachings of D1 and D2 in order to construct vectors having the technical features according to claims 9-12. Therefore, the subject-matter of claims 9-12 is considered to lack an inventive step.

The subject-matter according to claims 13-15 is novel. D3 is considered to represent the closest prior art. D3 describes E. coli strains with a mutation in the araD gene. The inactivation of araD blocks the utilisation of L-arabinose as a carbon source and leads to the accumulation of a toxic intermediate (see page 303, column 2, paragraph 3 and table 1).

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The subject-matter of claims 13-15 differs from what is disclosed in D3 in that other strains of E. coli are mutated in the araD gene. The E. coli strains which are used are the commercially available AG1, JM109 and DH5alpha-T1. It is uncertain what technical effect is achieved due to this difference.

In view of what is known from D3, and in the absence of an unexpected technical effect due to the above mentioned difference, it is considered as an obvious embodiment to the person skilled in the art to construct mutated strains according to claims 13-15. The subject-matter of claims 13-15 is therefore considered to lack an inventive step.

However, claims relating to the use of vectors and mutated strains as described in present claims 8-15 for the construction of a selection system according to the invention would be regarded as novel and inventive.